10/593958

March

- 1/12 - MP6 Rec'd PCT/PTO 22 SEP 2006

SEQUENCE LISTING

<110> BORGFORD, Thor BRAUN, Curtis PURAC, Admir STOLL, Dominik GLYCOSYLATION VARIANTS OF RICIN-LIKE PROTEINS <130> 10447-50 <140> <141> <150> US 60/555,678 <151> 2004-03-24 <160> <170> PatentIn version 3.3 <210> <211> 530 <212> PRT <213> Artificial Sequence <220> <223> TST10088 Protein Sequence Glu Ala Glu Ala Ile Phe Pro Lys Gln Tyr Pro Ile Ile Gln Phe Thr 5 10 Thr Ala Gly Ala Thr Val Gln Ser Tyr Thr Asn Phe Ile Arg Ala Val 20 25 Arg Gly Arg Leu Thr Thr Gly Ala Asp Val Arg His Glu Ile Pro Val 35 45 Leu Pro Asn Arg Val Gly Leu Pro Ile Asn Gln Arg Phe Ile Leu Val 50 Glu Leu Ser Asn His Ala Glu Leu Ser Val Thr Leu Ala Leu Asp Val 65 70 75 80 Thr Asn Ala Tyr Val Val Gly Tyr Arg Ala Gly Asn Ser Ala Tyr Phe 85 Phe His Pro Asp Asn Gln Glu Asp Ala Glu Ala Ile Thr His Leu Phe 100 Thr Asp Val Gln Asn Arg Tyr Thr Phe Ala Phe Gly Gly Asn Tyr Asp 115 120 125

Arg Leu Glu Gln Leu Ala Gly Asn Leu Arg Glu Asn Ile Glu Leu Gly 135 Asn Gly Pro Leu Glu Glu Ala Ile Ser Ala Leu Tyr Tyr Tyr Ser Thr Gly Gly Thr Gln Leu Pro Thr Leu Ala Arg Ser Phe Ile Ile Cys Ile 165 170 Gln Met Ile Ser Glu Ala Ala Arg Phe Gln Tyr Ile Glu Gly Glu Met 185 Arg Thr Arg Ile Arg Tyr Asn Arg Arg Ser Ala Pro Asp Pro Ser Val 200 Ile Thr Leu Glu Asn Ser Trp Gly Arg Leu Ser Thr Ala Ile Gln Glu 210 215 220 Ser Asn Gln Gly Ala Phe Ala Ser Pro Ile Gln Leu Gln Arg Arg Asn 225 230 235 Gly Ser Lys Phe Ser Val Tyr Asp Val Ser Ile Leu Ile Pro Ile Ile 250 Ala Leu Met Val Tyr Arg Cys Ser Pro Gln Gly Ile Ala Gly Gln Cys Met Asp Pro Glu Pro Ile Val Arg Ile Val Gly Arg Asn Gly Leu Cys 280 Val Asp Val Arg Asp Gly Arg Phe His Asn Gly Asn Ala Ile Gln Leu Trp Pro Cys Lys Ser Asn Thr Asp Ala Asn Gln Leu Trp Thr Leu Lys Arg Asp Asn Thr Ile Arg Ser Asn Gly Lys Cys Leu Thr Thr Tyr Gly 325 330 Tyr Ser Pro Gly Val Tyr Val Met Ile Tyr Asp Cys Asn Thr Ala Ala 345 Thr Asp Ala Thr Arg Trp Gln Ile Trp Asp Asn Gly Thr Ile Ile Asn 360

Pro Arg Ser Ser Leu Val Leu Ala Ala Thr Ser Gly Asn Ser Gly Thr 370 375 380

Thr Leu Thr Val Gln Thr Asn Ile Tyr Ala Val Ser Gln Gly Trp Leu 385 390 395 400

Pro Thr Gln Asn Thr Gln Pro Phe Val Thr Thr Ile Val Gly Leu Tyr 405 410 415

Gly Leu Cys Leu Gln Ala Asn Ser Gly Gln Val Trp Ile Glu Asp Cys 420 425 430

Ser Ser Glu Lys Ala Glu Gln Gln Trp Ala Leu Tyr Ala Asp Gly Ser 435 440 445

Ile Arg Pro Gln Gln Asn Arg Asp Asn Cys Leu Thr Ser Asp Ser Asn 450 455 460

Ile Arg Glu Thr Val Val Lys Ile Leu Ser Cys Gly Pro Ala Ser Ser 465 470 475 480

Gly Gln Arg Trp Met Phe Lys Asn Asp Gly Thr Ile Leu Asn Leu Tyr 485 490 495

Ser Gly Leu Val Leu Asp Val Arg Ala Ser Asp Pro Ser Leu Lys Gln 500 505 510

Ile Ile Leu Tyr Pro Leu His Gly Asp Pro Asn Gln Ile Trp Leu Pro 515 520 525

Leu Phe 530

<210> 2

<211> 530

<212> PRT

<213> Artificial Sequence

<220>

<223> TST10092 Protein Sequence

<400> 2

Glu Ala Glu Ala Ile Phe Pro Lys Gln Tyr Pro Ile Ile Gln Phe Thr
1 5 10 15

Thr Ala Gly Ala Thr Val Gln Ser Tyr Thr Asn Phe Ile Arg Ala Val 20 25 30 Arg Gly Arg Leu Thr Thr Gly Ala Asp Val Arg His Glu Ile Pro Val Leu Pro Asn Arg Val Gly Leu Pro Ile Asn Gln Arg Phe Ile Leu Val Glu Leu Ser Asn His Ala Glu Leu Ser Val Thr Leu Ala Leu Asp Val 70 Thr Asn Ala Tyr Val Val Gly Tyr Arg Ala Gly Asn Ser Ala Tyr Phe Phe His Pro Asp Asn Gln Glu Asp Ala Glu Ala Ile Thr His Leu Phe Thr Asp Val Gln Asn Arg Tyr Thr Phe Ala Phe Gly Gly Asn Tyr Asp 115 120 Arg Leu Glu Gln Leu Ala Gly Asn Leu Arg Glu Asn Ile Glu Leu Gly 135 Asn Gly Pro Leu Glu Glu Ala Ile Ser Ala Leu Tyr Tyr Tyr Ser Thr 155 Gly Gly Thr Gln Leu Pro Thr Leu Ala Arg Ser Phe Ile Ile Cys Ile 170 Gln Met Ile Ser Glu Ala Ala Arg Phe Gln Tyr Ile Glu Gly Glu Met Arg Thr Arg Ile Arg Tyr Asn Arg Arg Ser Ala Pro Asp Pro Ser Val Ile Thr Leu Glu Asn Ser Trp Gly Arg Leu Ser Thr Ala Ile Gln Glu 215 220 Ser Asn Gln Gly Ala Phe Ala Ser Pro Ile Gln Leu Gln Arg Arg Asn 230 235 Gly Ser Lys Phe Ser Val Tyr Asp Val Ser Ile Leu Ile Pro Ile Ile 250

Ala Leu Met Val Tyr Arg Cys Ser Pro Gln Gly Ile Ala Gly Gln Cys 260 265 270 Met Asp Pro Glu Pro Ile Val Arg Ile Val Gly Arg Asn Gly Leu Cys 275 280 285

: :

- Val Asp Val Arg Asp Gly Arg Phe His Asn Gly Asn Ala Ile Gln Leu 290 295 300
- Trp Pro Cys Lys Ser Asn Thr Asp Ala Asn Gln Leu Trp Thr Leu Lys 305 310 315 320
- Arg Asp Asn Thr Ile Arg Ser Asn Gly Lys Cys Leu Thr Thr Tyr Gly 325 330 335
- Tyr Ser Pro Gly Val Tyr Val Met Ile Tyr Asp Cys Asn Thr Ala Ala 340 345 350
- Thr Asp Ala Thr Arg Trp Gln Ile Trp Asp Asn Gly Thr Ile Ile Asn 355 360 365
- Pro Arg Ser Ser Leu Val Leu Ala Ala Thr Ser Gly Asn Ser Gly Thr 370 375 380
- Thr Leu Thr Val Gln Thr Asn Ile Tyr Ala Val Ser Gln Gly Trp Leu 385 390 395 400
- Pro Thr Asn Asn Thr Gln Pro Phe Val Thr Thr Ile Val Gly Leu Tyr 405 410 415
- Gly Leu Cys Leu Gln Ala Asn Ser Gly Gln Val Trp Ile Glu Asp Cys 420 425 430
- Ser Ser Glu Lys Ala Glu Gln Gln Trp Ala Leu Tyr Ala Asp Gly Ser 435 440 445
- Ile Arg Pro Gln Gln Asn Arg Asp Asn Cys Leu Thr Ser Asp Ser Asn 450 455 460
- Ile Arg Glu Thr Val Val Lys Ile Leu Ser Cys Gly Pro Ala Ser Ser 465 470 475 480
- Gly Gln Arg Trp Met Phe Lys Asn Asp Gly Thr Ile Leu Asn Leu Tyr 485 490 495
- Ser Gly Leu Val Leu Asp Val Arg Ala Ser Asp Pro Ser Leu Lys Gln 500 505 510

Ile Ile Leu Tyr Pro Leu His Gly Asp Pro Asn Gln Ile Trp Leu Pro 515 520 525

Leu Phe 530

<210> 3

<211> 531

<212> PRT

<213> Artificial Sequence

:

<220>

<223> TST10147 Protein Sequence

<400> 3

Thr Ala Gly Ala Thr Val Gln Ser Tyr Thr Asn Phe Ile Arg Ala Val 20 25 30

Arg Gly Arg Leu Thr Thr Gly Ala Asp Val Arg His Glu Ile Pro Val 35 40 45

Leu Pro Asn Arg Val Gly Leu Pro Ile Asn Gln Arg Phe Ile Leu Val 50 55 60

Glu Leu Ser Asn His Ala Glu Leu Ser Val Thr Leu Ala Leu Asp Val 65 70 75 80

Thr Asn Ala Tyr Val Val Gly Tyr Arg Ala Gly Asn Ser Ala Tyr Phe 85 90 95

Phe His Pro Asp Asn Gln Glu Asp Ala Glu Ala Ile Thr His Leu Phe 100 105 110

Thr Asp Val Gln Asn Arg Tyr Thr Phe Ala Phe Gly Gly Asn Tyr Asp 115 120 125

Arg Leu Glu Gln Leu Ala Gly Asn Leu Arg Glu Asn Ile Glu Leu Gly 130 135 140

Asn Gly Pro Leu Glu Glu Ala Ile Ser Ala Leu Tyr Tyr Tyr Ser Thr 145 150 155 160

Gly Gly Thr Gln Leu Pro Thr Leu Ala Arg Ser Phe Ile Ile Cys Ile 165 170 175

- Gln Met Ile Ser Glu Ala Ala Arg Phe Gln Tyr Ile Glu Gly Glu Met
 180 185 190

 Arg Thr Arg Ile Arg Tyr Asn Arg Arg Ser Ala Pro Asp Pro Ser Val
- Ile Thr Leu Glu Asn Ser Trp Gly Arg Leu Ser Thr Ala Ile Gln Glu 210 215 220
- Ser Asn Gln Gly Ala Phe Ala Ser Pro Ile Gln Leu Gln Arg Arg Asn 225 230 235 240
- Gly Ser Lys Phe Ser Val Tyr Asp Val Ser Ile Leu Ile Pro Ile Ile 245 250 255
- Ala Leu Met Val Tyr Arg Cys Gly Ser Pro Gln Gly Ile Ala Gly Gln 260 265 270
- Cys Met Asp Pro Glu Pro Ile Val Arg Ile Val Gly Arg Asn Gly Leu 275 280 285
- Cys Val Asp Val Arg Asp Gly Arg Phe His Asn Gly Asn Ala Ile Gln 290 295 300
- Leu Trp Pro Cys Lys Ser Asn Thr Asp Ala Asn Gln Leu Trp Thr Leu 305 310 315 320
- Lys Arg Asp Asn Thr Ile Arg Ser Asn Gly Lys Cys Leu Thr Thr Tyr 325 330 335
- Gly Tyr Ser Pro Gly Val Tyr Val Met Ile Tyr Asp Cys Asn Thr Ala 340 345 350
- Ala Thr Asp Ala Thr Arg Trp Gln Ile Trp Asp Asn Gly Thr Ile Ile 355 360 365
- Asn Pro Arg Ser Ser Leu Val Leu Ala Ala Thr Ser Gly Asn Ser Gly 370 375 380
- Thr Thr Leu Thr Val Gln Thr Asn Ile Tyr Ala Val Ser Gln Gly Trp 385 390 395 400
- Leu Pro Thr Gln Asn Thr Gln Pro Phe Val Thr Thr Ile Val Gly Leu 405 410 415

Tyr Gly Leu Cys Leu Gln Ala Asn Ser Gly Gln Val Trp Ile Glu Asp 420 425 430 Cys Ser Ser Glu Lys Ala Glu Gln Gln Trp Ala Leu Tyr Ala Asp Gly 440 Ser Ile Arg Pro Gln Gln Asn Arg Asp Asn Cys Leu Thr Ser Asp Ser 455 Asn Ile Arg Glu Thr Val Val Lys Ile Leu Ser Cys Gly Pro Ala Ser 465 470 475 480 Ser Gly Gln Arg Trp Met Phe Lys Asn Asp Gly Thr Ile Leu Asn Leu 485 490 Tyr Ser Gly Leu Val Leu Asp Val Arg Ala Ser Asp Pro Ser Leu Lys 500 505 Gln Ile Ile Leu Tyr Pro Leu His Gly Asp Pro Asn Gln Ile Trp Leu 515 520 525 Pro Leu Phe 530 <210> <211> 1707 <212> DNA <213> Artificial Sequence <220> <223> TST10088 DNA Insert Sequence <220> <221> prim_transcript <222> (1)..(117)<400> 4 atgaaaccgg gaggaaatac tattgtaata tgggtgtatg cagtggcaac atggctttgt 60 tttggatcca cctcagggtg gtctttcaca ttagaggata acaacctcga gaaaagagag 120 gctgaagcta tattccccaa acaataccca attatacagt ttaccacagc gggtgccact 180 gtgcaaagct acacaaactt tatcagagct gttcgcggtc gtttaacaac tggagctgat 240 gtgagacatg aaataccagt gttgccaaac agagttggtt tgcctataaa ccaacggttt 300 attttagttg aacteteaaa teatgeagag etttetgtta eattageget ggatgteace 360 aatgcatatg tggtcggcta ccgtgctgga aatagcgcat atttctttca tcctgacaat 420 caggaagatg cagaagcaat cactcatctt ttcactgatg ttcaaaaatcg atatacattc 480 gcctttggtg gtaattatga tagacttgaa caacttgctg gtaatctgag agaaaatatc 540 gagttgggaa atggtccact agaggaggct atctcagcgc tttattatta cagtactggt 600 ggcactcage ttecaactet ggetegttee tttataattt geateeaaat gattteagaa 660 gcagcaagat tccaatatat tgagggagaa atgcgcacga gaattaggta caaccggaga 720 tctgcaccag atcctagcgt aattacactt gagaatagtt gggggagact ttccactgca 780 attcaagagt ctaaccaagg agcctttgct agtccaattc aactgcagag acgtaatggt 840 tocaaattoa gtgtgtacga tgtgagtata ttaatcocta tcatagotot catggtgtat 900 agatgctctc cgcaaggaat tgcagggcag tgtatggatc ctgagcccat agtgcgtatc 960 1020 gtaggtcgaa atggtctatg tgttgatgtt agggatggaa gattccacaa cggaaacgca 1080 atacagttgt ggccatgcaa gtctaataca gatgcaaatc agctctggac tttgaaaaga gacaatacta ttcgatctaa tggaaagtgt ttaactactt acgggtacag tccgggagtc 1140 tatgtgatga tctatgattg caatactgct gcaactgatg ccacccgctg gcaaatatgg 1200 gataatggaa ccatcataaa tcccagatct agtctagttt tagcagcgac atcagggaac 1260 agtggtacca cacttacagt gcaaaccaac atttatgccg ttagtcaagg ttggcttcct 1320 actcagaata cacaaccttt tgtgacaacc attgttgggc tatatggtct gtgcttgcaa 1380 gcaaatagtg gacaagtatg gatagaggac tgtagcagtg aaaaggctga acaacagtgg 1440 gctctttatg cagatggttc aatacgtcct cagcaaaacc gagataattg ccttacaagt 1500 gattctaata tacgggaaac agttgtcaag atcctctctt gtggccctgc atcctctggc 1560 caacgatgga tgttcaagaa tgatggaacc attttaaatt tgtatagtgg gttggtgtta 1620 1680 gatgtgaggg catcagatcc gagccttaaa caaatcattc tttaccctct ccatggtgac ccaaaccaaa tatggttacc attattt 1707

```
<210> 5
<211> 1707
<212> DNA
<213> Artificial Sequence
<220>
<223> TST10092 DNA Insert Sequence
```

<220>
<221> prim_transcript
<222> (1)..(117)

<400> 5
atgaaaccgg gaggaaatac tattgtaata tgggtgtatg cagtggcaac atggctttgt 60
tttggatcca cctcagggtg gtctttcaca ttagaggata acaacctcga gaaaagagag 120

180 gctgaagcta tattccccaa acaataccca attatacagt ttaccacagc gggtgccact gtgcaaagct acacaaactt tatcagagct gttcgcggtc gtttaacaac tggagctgat 240 gtgagacatg aaataccagt gttgccaaac agagttggtt tgcctataaa ccaacggttt 300 360 attttagttg aactctcaaa tcatgcagag ctttctgtta cattagcgct ggatgtcacc aatgcatatg tggtcggcta ccgtgctgga aatagcgcat atttctttca tcctgacaat 420 caggaagatg cagaagcaat cactcatctt ttcactgatg ttcaaaatcg atatacattc 480 gcctttggtg gtaattatga tagacttgaa caacttgctg gtaatctgag agaaaatatc 540 600 gagttgggaa atggtccact agaggaggct atctcagcgc tttattatta cagtactggt ggcactcage ttccaactct ggctcgttcc tttataattt gcatccaaat gatttcagaa 660 720 gcagcaagat tccaatatat tgagggagaa atgcgcacga gaattaggta caaccggaga tctgcaccag atcctagcgt aattacactt gagaatagtt gggggagact ttccactgca 780 attcaagagt ctaaccaagg agcctttgct agtccaattc aactgcagag acgtaatggt 840 tccaaattca gtgtgtacga tgtgagtata ttaatcccta tcatagctct catggtgtat 900 agatgetete egcaaggaat tgeagggeag tgtatggate etgageeeat agtgegtate 960 1020 gtaggtcgaa atggtctatg tgttgatgtt agggatggaa gattccacaa cggaaacgca 1080 atacagttgt ggccatgcaa gtctaataca gatgcaaatc agctctggac tttgaaaaga 1140 gacaatacta ttcgatctaa tggaaagtgt ttaactactt acgggtacag tccgggagtc tatgtgatga totatgattg caatactgct gcaactgatg ccacccgctg gcaaatatgg 1200 gataatggaa ccatcataaa tcccagatct agtctagttt tagcagcgac atcagggaac 1260 agtggtacca cacttacagt gcaaaccaac atttatgccg ttagtcaagg ttggcttcct 1320 1380 actaataata cacaaccttt tgtgacaacc attgttgggc tatatggtct gtgcttgcaa 1440 gcaaatagtg gacaagtatg gatagaggac tgtagcagtg aaaaggctga acaacagtgg 1500 gctctttatg cagatggttc aatacgtcct cagcaaaacc gagataattg ccttacaagt gattctaata tacgggaaac agttgtcaag atcctctctt gtggccctgc atcctctggc 1560 caacgatgga tgttcaagaa tgatggaacc attttaaatt tgtatagtgg gttggtgtta 1620 gatgtgaggg catcagatcc gagccttaaa caaatcattc tttaccctct ccatggtgac 1680 1707 ccaaaccaaa tatggttacc attattt

<210> 6

<211> 1710

<212> DNA

<213> Artificial Sequence

<220> <223> TST10147 DNA Insert Sequence

<220>
<221> prim_transcript
<222> (1)..(117)

<400> atgaaaccgg gaggaaatac tattgtaata tgggtgtatg cagtggcaac atggctttgt 60 tttggatcca cctcagggtg gtctttcaca ttagaggata acaacctcga gaaaagagag 120 gctgaagcta tattccccaa acaataccca attatacagt ttaccacagc gggtgccact 180 gtgcaaagct acacaaactt tatcagagct gttcgcggtc gtttaacaac tggagctgat 240 300 gtgagacatg aaataccagt gttgccaaac agagttggtt tgcctataaa ccaacggttt attttagttg aactctcaaa tcatgcagag ctttctgtta cattagcgct ggatgtcacc 360 aatgcatatg tggtcggcta ccgtgctgga aatagcgcat atttctttca tcctgacaat 420 caggaagatg cagaagcaat cactcatctt ttcactgatg ttcaaaatcg atatacattc 480 gcctttggtg gtaattatga tagacttgaa caacttgctg gtaatctgag agaaaatatc 540 gagttgggaa atggtccact agaggaggct atctcagcgc tttattatta cagtactggt 600 ggcactcage ttecaactet ggctegttee tttataattt gcatecaaat gattteagaa 660 gcagcaagat tccaatatat tgagggagaa atgcgcacga gaattaggta caaccggaga 720 tctgcaccag atcctagcgt aattacactt gagaatagtt gggggagact ttccactgca 780 attcaagagt ctaaccaagg agcctttgct agtccaattc aactgcagag acgtaatggt 840 tccaaattca gtgtgtacga tgtgagtata ttaatcccta tcatagctct catggtgtat 900 agatgcggtt ctccgcaagg aattgcaggg cagtgtatgg atcctgagcc catagtgcgt 960 1020 atcgtaggtc gaaatggtct atgtgttgat gttagggatg gaagattcca caacggaaac 1080 gcaatacagt tgtggccatg caagtctaat acagatgcaa atcagctctg gactttgaaa agagacaata ctattcgatc taatggaaag tgtttaacta cttacgggta cagtccggga 1140 gtctatgtga tgatctatga ttgcaatact gctgcaactg atgccacccg ctggcaaata 1200 1260 aacagtggta ccacacttac agtgcaaacc aacatttatg ccgttagtca aggttggctt 1320 cctactcaga atacacaacc ttttgtgaca accattgttg ggctatatgg tctgtgcttg 1380 caagcaaata gtggacaagt atggatagag gactgtagca gtgaaaaggc tgaacaacag 1440 tgggctcttt atgcagatgg ttcaatacgt cctcagcaaa accgagataa ttgccttaca 1500 agtgatteta atataeggga aacagttgte aagateetet ettgtggeee tgeateetet 1560 ggccaacgat ggatgttcaa gaatgatgga accattttaa atttgtatag tgggttggtg 1620 ttagatgtga gggcatcaga tccgagcctt aaacaaatca ttctttaccc tctccatggt 1680 4.